

SEQUENCE LISTING

<110> SHERMAN, LINDA A. LUSTGARTEN, JOSEPH <120> RECOMBINANT CONSTRUCTS ENCODING T CELL RECEPTORS SPECIFIC FOR HUMAN HLA-RESTRICTED TUMOR ANTIGENS <130> 48340/55793 <140> 08/812,393 <141> 1997-03-05 <150> 60/012,845 <151> 1996-03-05 <160> 64 <170> PatentIn Ver. 2.1 <210> 1 <211> 1350 <212> DNA <213> Artificial Sequence <220> <221> CDS <222> (1)..(1332) <220> <223> Description of Artificial Sequence: Synthetic single chain TCR derivative nucleotide sequence <400> 1 ctc gag atg cag agg aac ctg gga gct gtg ctg ggg att ctg tgg gtg Leu Glu Met Gln Arg Asn Leu Gly Ala Val Leu Gly Ile Leu Trp Val 48 10 cag att tgc tgg ctg aaa gaa cag caa gtg cag cag agt ccc gca tcc 96 Gln Ile Cys Trp Leu Lys Glu Gln Gln Val Gln Gln Ser Pro Ala Ser ttg gtt ctg cag gag ggg gag aac gca gag ctc cag tgt agc ttt tcc Leu Val Leu Gln Glu Gly Glu Asn Ala Glu Leu Gln Cys Ser Phe Ser atc ttt aca aac cag gtg cag tgg ttt tac caa cgt cct ggg gga aga Ile Phe Thr Asn Gln Val Gln Trp Phe Tyr Gln Arg Pro Gly Gly Arg ctc gtc agc ctg ttg tac aat cct tct ggg aca aag cag agt ggg aga 240 Leu Val Ser Leu Leu Tyr Asn Pro Ser Gly Thr Lys Gln Ser Gly Arg 70 ctg aca tcc aca aca gtc att aaa gaa cgt cgc agc tct ttg cac att Leu Thr Ser Thr Thr Val Ile Lys Glu Arg Arg Ser Ser Leu His Ile 85 90

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Pro	Val	Phe 275	Leu	Pro	Ala	Lys	280	Thr	Thr	Thr		285	Pro	Arg	Pro	864
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tct Ser 305	Ser	tct Ser	aga Arg	gat JAsp	2 ccc 2 Pro 310	Lys	cto Lev	tgo Cys	tac Tyr	Let 315	ı Lev	gat Asp	gga Gly	a ato	Leu 320	960

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Glr	ı Ile	е Сув	Trp		Lys	Glu	ı Glr	ı Glr 25		. Glr	Glr	Ser	Pro		Ser	
Let	ı Val	. Let		Glu	ı Gly	Glu	a Asr 4(a Glu	ı Lev	Glr	Cys		Phe	e Ser	
Ile	e Phe		Asr	Glr	ı Val	. Glr 55		Phe	е Туг	Glr	Arg		Gly	/ Gly	Arg	

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Leu Thr Ser Thr Thr Val Ile Lys Glu Arg Arg Ser Ser Leu His Ile 85 90 95

Ser Ser Ser Gln Ile Thr Asp Ser Gly Thr Tyr Leu Cys Ala Ser Asn 100 105 110

Ser Gly Gly Ser Asn Ala Lys Leu Thr Phe Gly Lys Gly Thr Lys Leu 115 120 125

Ser Val Lys Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Gly Gly 130 135 140

Gly Gly Ser Glu Ala Ala Val Thr Gln Ser Pro Arg Asn Lys Val Ala 145 150 155 160

Val Thr Gly Gly Lys Val Thr Leu Ser Cys Asn Gln Thr Asn Asn His

Asn Asn Met Tyr Trp Tyr Arg Gln Asp Thr Gly His Gly Leu Arg Leu 180 185 190

Ile His Tyr Ser Tyr Gly Ala Gly Ser Thr Glu Lys Gly Asp Ile Pro 195 200 205

Asp Gly Tyr Lys Ala Ser Arg Pro Ser Gln Glu Asn Phe Ser Leu Ile 210 215 220

Leu Glu Leu Ala Thr Pro Ser Gln Thr Ser Val Tyr Phe Cys Ala Ser 225 230 235 240

Gly Glu Thr Gly Thr Asn Glu Arg Leu Phe Phe Gly His Gly Thr Lys 245 250 255

Leu Ser Val Leu Thr Ser Asn Ser Ile Met Tyr Phe Ser His Phe Val 260 265 270

Pro Val Phe Leu Pro Ala Lys Pro Thr Thr Thr Pro Ala Pro Arg Pro 275 280 285

Pro Thr Pro Ala Pro Thr Ile Ala Ser Gln Pro Leu Ser Leu Arg Pro 290 295 300

Ser Ser Ser Arg Asp Pro Lys Leu Cys Tyr Leu Leu Asp Gly Ile Leu 305 310 320

Phe Ile Tyr Gly Val Ile Leu Thr Ala Leu Phe Leu Arg Val Lys Phe 325 330 335

Ser Arg Ser Ala Asp Ala Pro Ala Tyr Gln Gln Gly Gln Asn Gln Leu 340 345 350

Tyr Asn Glu Leu Asn Leu Gly Arg Arg Glu Glu Tyr Asp Val Leu Asp 355 360 365

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Glu	Ala	Tyr	Ser	Glu 405	Ile	Gly	Met	Lys	Gly 410	Glu	Arg	Arg	Arg	Gly 415	Lys	
Gly	His	Asp	Gly 420	Leu	Tyr	Gln	Gly	Leu 425	Ser	Thr	Ala	Thr	Lys 430	Asp	Thr	
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												aaa Lys				240
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Tyr Met Tyr Trp Tyr Arg Gln Asp Thr Gly His Gly Leu Arg Leu Ile 50 55 60

His Tyr Ser Tyr Val Ala Asp Ser Thr Glu Lys Gly Asp Ile Pro Asp 65 70 75 80

Gly Tyr Lys Ala Ser Arg Pro Ser Gln Glu Asn Phe Ser Leu Ile Leu 85 90 95

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- Ser Gly Gly Ser Asn Ala Lys Leu Thr Phe Gly Lys Gly Thr Lys Leu 115 120 125
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- Val Thr Gly Gly Lys Val Thr Leu Ser Cys Asn Gln Thr Asn Asn His
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- Ile His Tyr Ser Tyr Gly Ala Gly Ser Thr Glu Lys Gly Asp Ile Pro 195 200 205
- Asp Gly Tyr Lys Ala Ser Arg Pro Ser Gln Glu Asn Phe Ser Leu Ile 210 215 220
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Thr Gly Gly Lys Val Thr Leu Ser Cys His Gln Thr Asn Asn His Asp 35 40 45

Tyr Met Tyr Trp Tyr Arg Gln Asp Thr Gly His Gly Leu Arg Leu Ile 50 60

His Tyr Ser Tyr Val Ala Asp Ser Thr Glu Lys Gly Asp Ile Pro Asp 65 70 75 80

Gly Tyr Lys Ala Ser Arg Pro Ser Gln Glu Asn Phe Ser Leu Ile Leu 85 90 95

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<223> Description of Artificial Sequence: Synthetic
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